

Table S2. Phase residuals in resolution shells for *p6* averaged Fourier terms of frozen hydrated CsxA crystals

Resolution shell (Å)	Number of independent reflections	Mean value of phase error against symmetry imposed phase of 0° or 180° (45° is expected for random phases)	Standard error (°)
200-15	25	11.0	4.0
15-12	14	31.9	6.9
12-10	13	25.3	4.4
10-9	18	25.9	5.9